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ANTIGEN 1

10	20	30	40	50	60	70	80	90	100
GAATTGGCACGGAAATTACCATCTGATAGATCAAATTACTACATCTTACTACATTAAATATGGAGGAAAACAGTCATGAGCAATCCACAA									
A R E L P S D R S N L L T S I F T T L N M E E K Q S M S N P Q									
5	10	15	20	25	30				
110	120	130	140	150	160	170	180	190	200
TCGAAAATAACGAATAACGAAATACCAACCAAAAGATTCTGGTTAAATGATAAAATATTGAAATGATTACAGATGAATTCAAAAAATTGACCTTAA									
S K N T N T S N T N H K D S G L N D K I F E M I T D E F K K L T F S									
35	40	45	50	55	60				
210	220	230	240	250	260	270	280	290	300
GCTTGTCCAAGAAATTAAATGATTCTGGTTCTTCAGCAATTAGCAAGTATTAGCAATTAGCAAGTAAATTCTGATGATACTTAAAGTCGCATTGTCAAGGA									
L S K E L N D S V S S A I S K Y L E P I E R D I H L L S R I C Q E									
5	70	75	80	85	90	95			
310	320	330	340	350	360	370	380	390	400
ATCGAGAAAGTCTGTGATAATTATGTTAAATCAATGAAATTCTAAATTGAAACTTCTAAATTGAAACAAATGTTAAGGAACACTTCTTACAAAGTACAAATGAGAAATTAAACAA									
S R S L L I I M L I S M K F L K L Q M L R N F L Q V Q M R N									
100	105	110	115	120	125				
410	420	430	440	450	460	470	480	490	500
GCATCGACACTTGTATTTCGAGGCTTGTGGCGAATCTAGAAGCGTTCTGTGAAAAGTGACTAAATTAAATAACATGGATAAACATTAAACTCGAATCC									
510	520	530	540	550	560	570	580	590	600
AATAGACAACTTACTCAAG TAGTAGCAGATTCTTACACCGATAACTCAATTACAATAAGAGCCCAAAACATAGCTTGTCAAATTGATGATGCCTTAAACCAA									
610	620	630	640	650	660	670	680	690	700
AGTAATGGAATAACCAACTAAAACGTCATTGCGAACTTAATGCAATTACAATAAGAGCCCAAAACATAGCTTGTCAAATTGATGATGCCTTAAACCAA									
710	720	730	740	750	760	770	780	790	800
AACATTACGATCGGCATTTCGGATAGCAATTCTGGATCAATCTAACTCTATCTCAATTGAGATAAGAGAGAAATCCAAGGGAGAATGTTTGTGAAAC									

FIG. 1A

□ 9 12 21 15 15 15 □ 10 15 □ 11 14 13

FIG. 1B

ANTIGEN 2

FIG. 2A

710 720 730 740 750 760 770 780 790 800  
 CCTCAAATGGTGAACATCAAGAATCTGGTTAGAGCAAGCCCTATTACCTCACCAAGAAACGAATCAAGTCAAATCATCCCTCTGTGACAGTCCAGA  
 810 820 830 840 850 860 870 880 890 900  
 TACTGGATCAGTTCAAATCTCCCTCTGTTACTATTCCAGAGACTGGATCAGACTCAGATCACGGGCCCTGTGACAATTCCAGAGACTGGATCAGTTCAA  
 910 920 930 940 950 960 970 980 990 1000  
 TCATCTCTGCTACTATACCAAGAACAGGATCCAGCTCAGATCACACTCTGCTACTTCTCCAGAAGAAGGATTGGACTCAGAACGTTACCAATCACTTCT  
 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100  
 ACAGAACAAACTCAAAGCCAGCTACATATCCTAACCAAGAAAATGAAAATCATAAATTCAGGAAGGTAATTTCGAGTTTAATAACACTAAATCTCCAA  
 1110 1120 1130 1140 1150 1160 1170 1180 1190 1200  
 TCAACCCCAATCTTCAAGCTGGCAATGTGAAAGTTATGGGAAAAGGATAAAATGGTTGATGGTGAGCAAGTAAATCACTAAAATGGACATTAT  
 1210 1220 1230 1240 1250 1260 1270 1280 1290 1300  
 GAAGATACTTCGAAAGAAATTAGAACAAAATGTAAGTATCTGCATTGATAAAATGGCCTTAGGCCATTTCAAATATCTAAATTGTCAACTCAAGTAA  
 1310 1320  
 AAAAAAAAAAACTCGAG

FIG. 2B

KD

45 -  
31 -  
21.5 -  
14.5 -



FIG. 3A

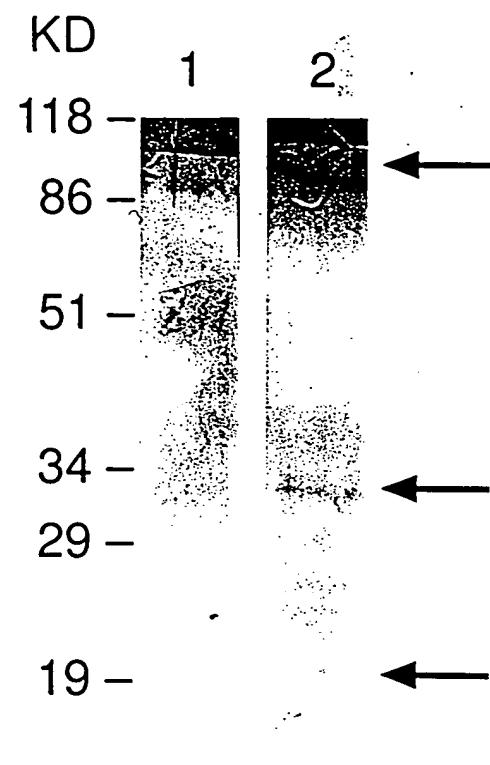


FIG. 3B

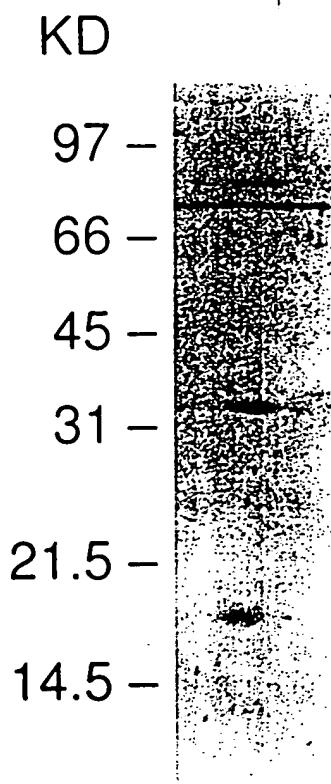


FIG. 4A

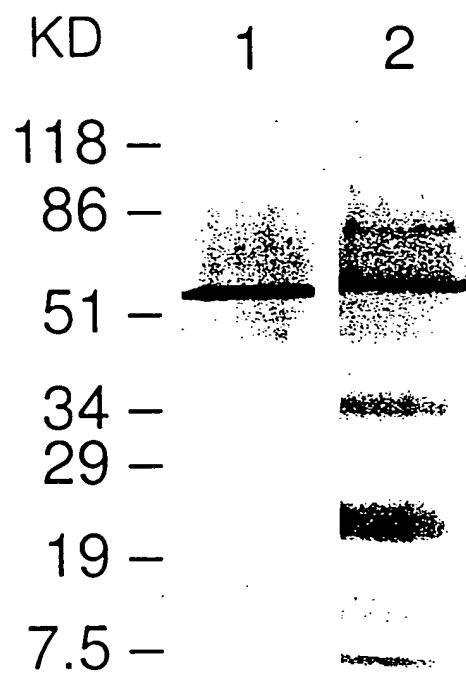


FIG. 4B

00000000000000000000000000000000

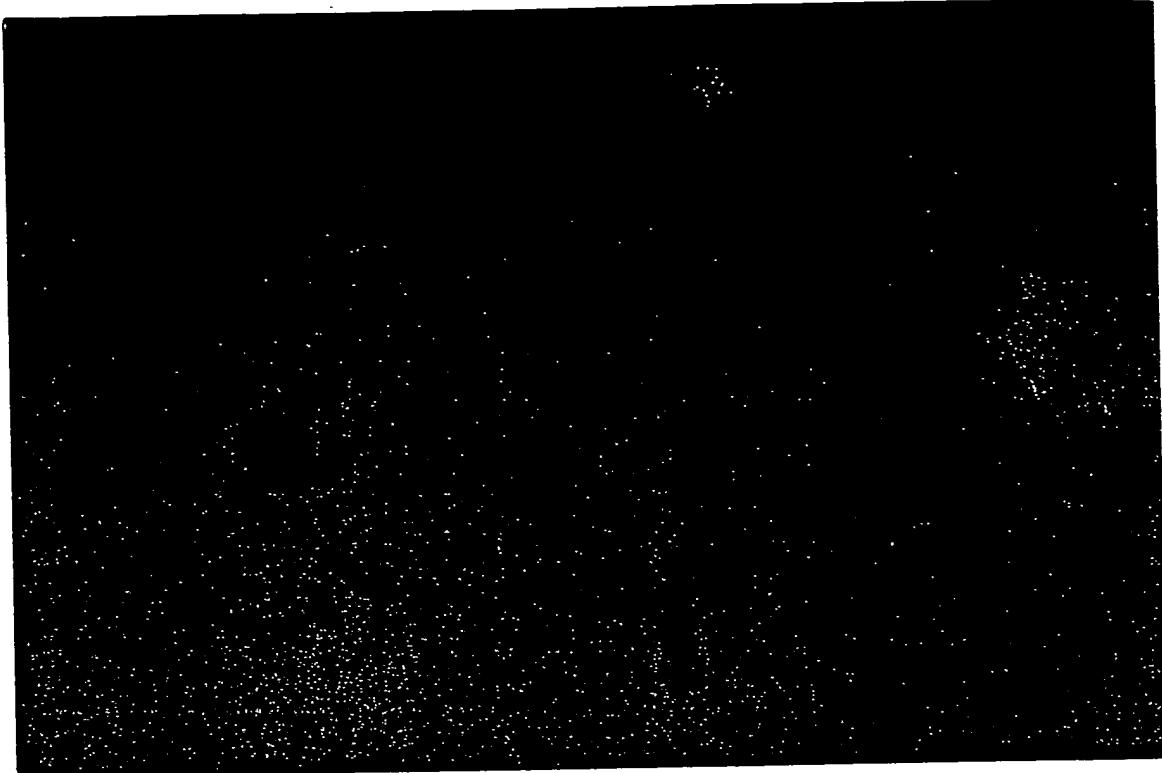


FIG. 5A

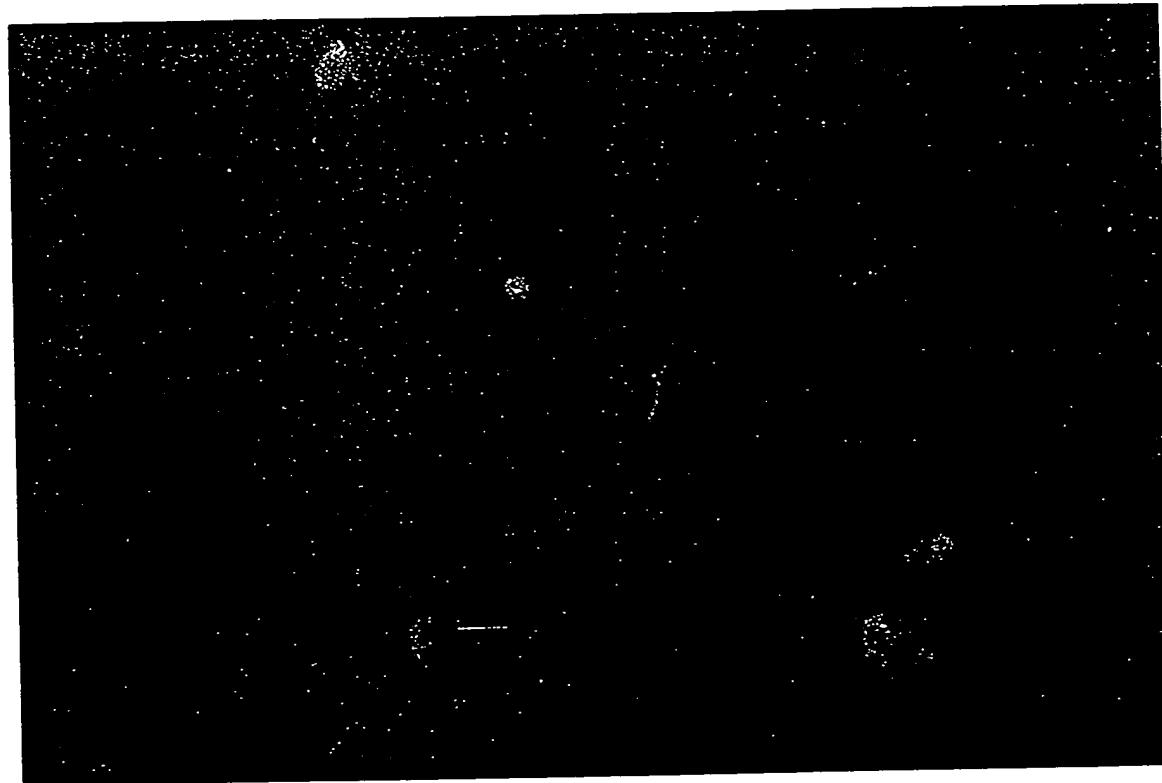


FIG. 5B

05291556 = 052900

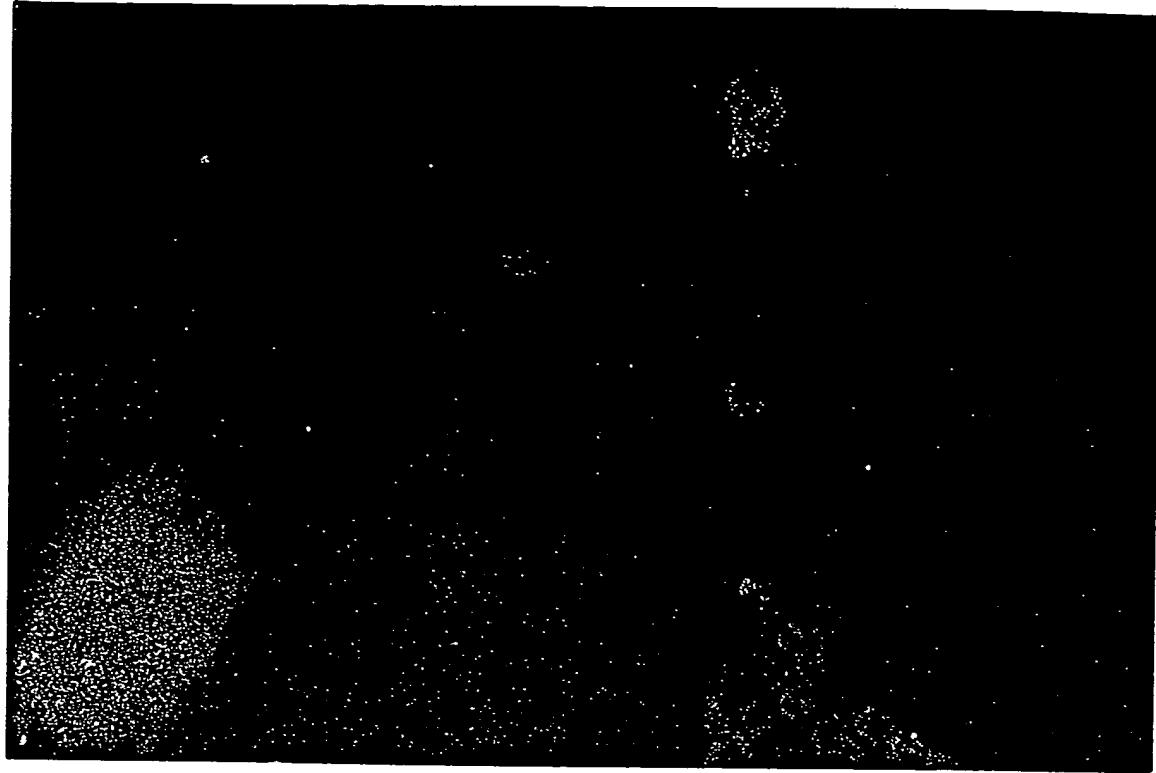


FIG 5C

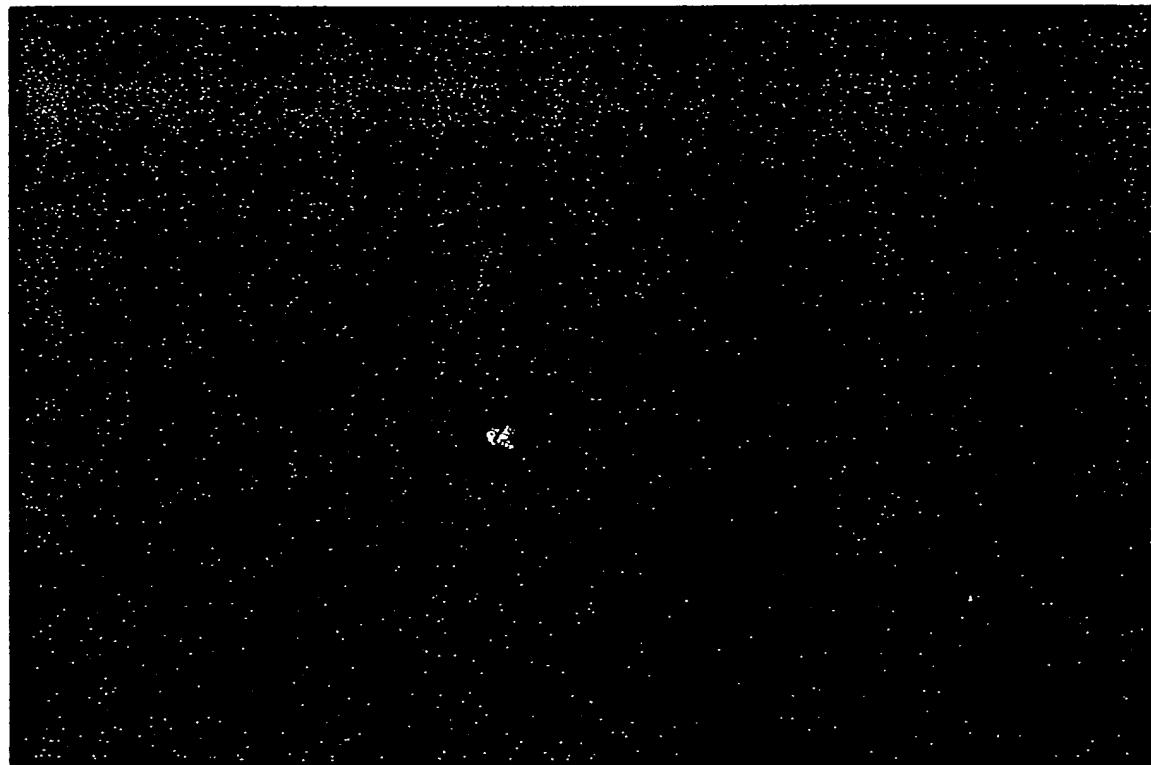


FIG 5D

1000 1000 1000 1000 1000



FIG 5E

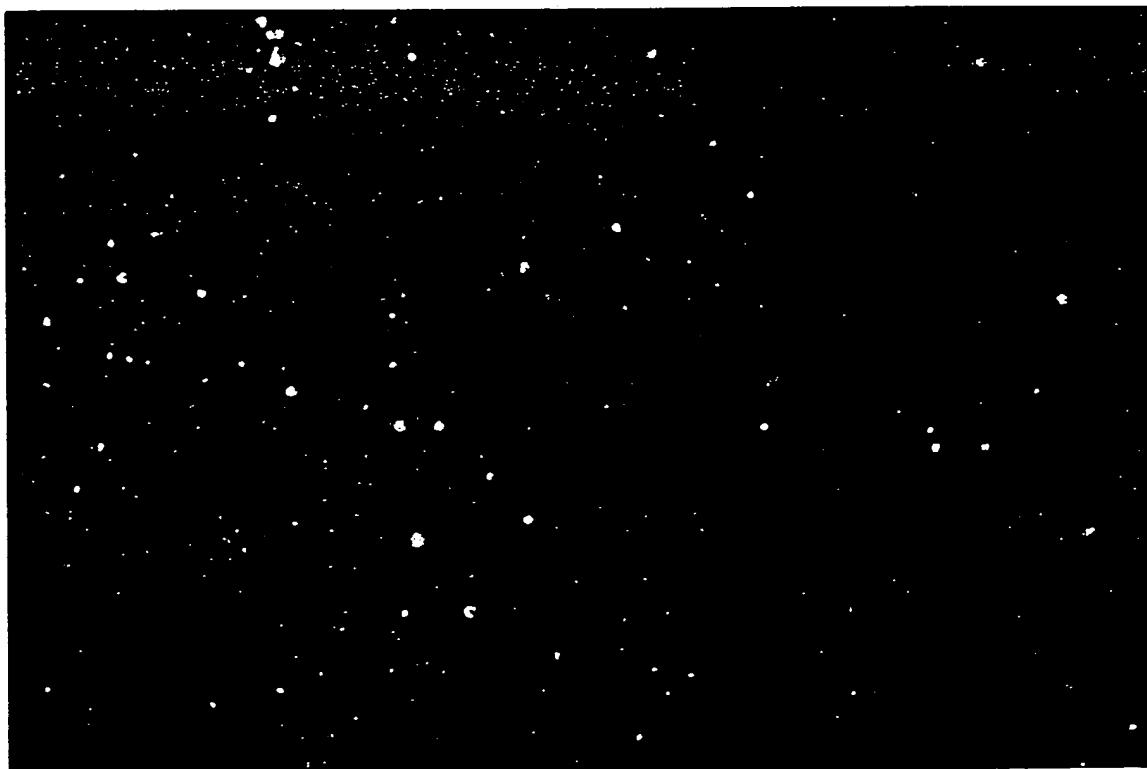


FIG 5F

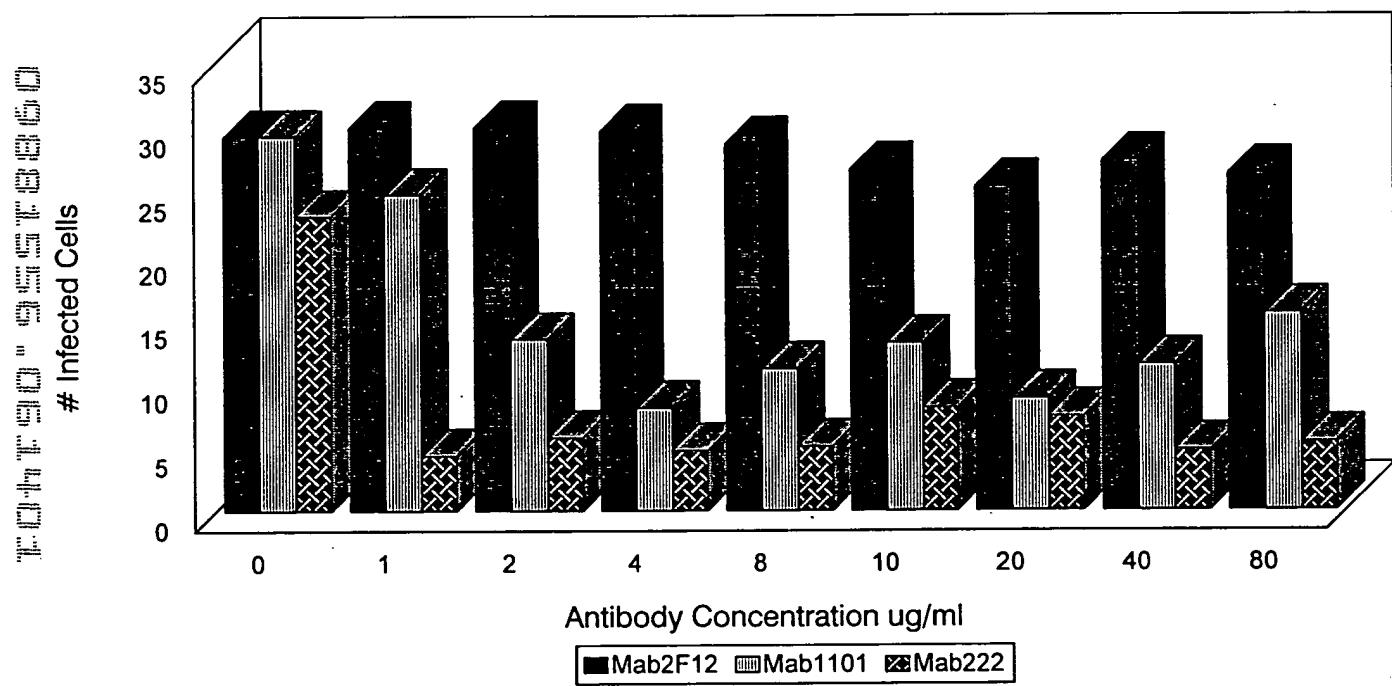


FIG. 6

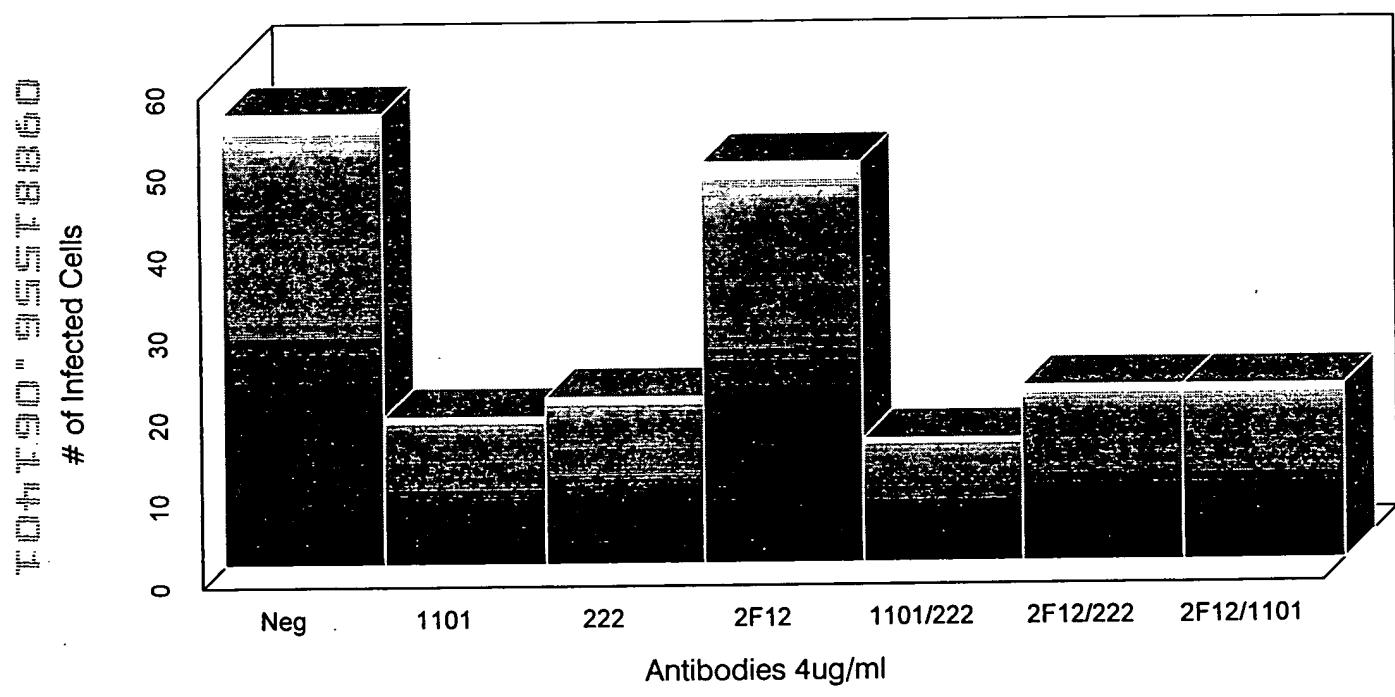


FIG. 7